

FIGURE 1A

387	AGA TCT CTC	396	CCT CAT ATA	405	GAT GTT TCT	414	TGG CGC TTG	423	GAA TAT CAG	432	ATA AAG
	R S L P H I T D V S W R L E Y Q I K										
441	ACC AAT CAA CTT	450	CAT AGG ATG	459	TAC AGA CCT	468	GCA TAT TTG	477	GTG ACC TTA	486	AGT GTA
	T N Q L H R M Y R P A Y L V T L S V										
495	CAG AAC ACT GAT	504	TCC CCA TCC	513	TAT CCA GAG	522	ATT AGT TTT	531	AGT TGC AGC	540	ATG GAA
	Q N T D S P S Y P E I S F S C S M E										
549	CAA TTA CAG GAC	558	TTG GTG GGG	567	AAA CTT AAA	576	GCT GCT TCG	585	AGC AAA	594	CTG GAA
	Q L Q D L V G K L K D A S K S L E R										
603	GCA ACT CAG TTG	612	TAA CTT GGG	621	GAA GTT AAC	630	GAT CCG CCC	639	GAG TGC AGA	648	GGA AAA
	A T Q L										
657	CCA GAA ACG CCT	666	TGC CTT CAG	675	CTG AAC CAC	684	CGT TTG TGC	693	GAG CTG GAT	702	GTC CTT
711	TTC AGT AGA	720	AAA GAA TTT	729	TCC TTT TGA	738	ATT TAT ACC	747	CAN CAA TTT	756	T T 3'

FIGURE 1B

1	MELSES	VQ	KG	FQ	ML	AD	PR	SF	DS	NA	F	T	L	L	R	A	A	F	Q	S	L	L	D	A	Q	2267574	
1	MELSES	VQ	RG	IT	QT	LA	DP	GS	FD	SN	A	F	A	L	L	R	A	A	F	Q	S	L	L	D	A	R	GI 265569
41	A	D	E	A	V	L	D	H	P	D	L	K	H	I	D	P	V	V	L	K	H	C	H	A	A	A	2267574
41	A	D	E	A	A	L	D	H	P	Y	L	K	Q	I	D	P	V	V	L	K	H	C	H	A	A	A	GI 265569
81	S	T	L	S	T	Y	L	E	D	C	K	F	D	R	E	R	I	E	L	F	C	T	E	Y	Q	N	2267574
81	S	T	L	S	T	Y	L	E	D	C	K	F	D	R	E	R	I	E	L	F	C	T	E	Y	Q	N	GI 265569
121	S	L	P	H	I	T	D	V	S	W	R	L	E	Y	Q	I	K	T	N	Q	L	H	R	M	Y	R	2267574
121	S	L	P	H	I	T	D	V	S	W	R	L	E	Y	Q	I	K	T	N	Q	L	H	K	M	Y	R	GI 265569
161	P	S	Y	P	E	I	S	F	S	C	S	M	E	Q	L	Q	D	L	V	G	K	L	K	D	A	S	2267574
161	Q	S	Y	P	E	I	N	F	S	C	N	M	E	Q	L	Q	D	L	V	G	K	L	K	D	A	S	GI 265569

FIGURE 2

16090" 31681800

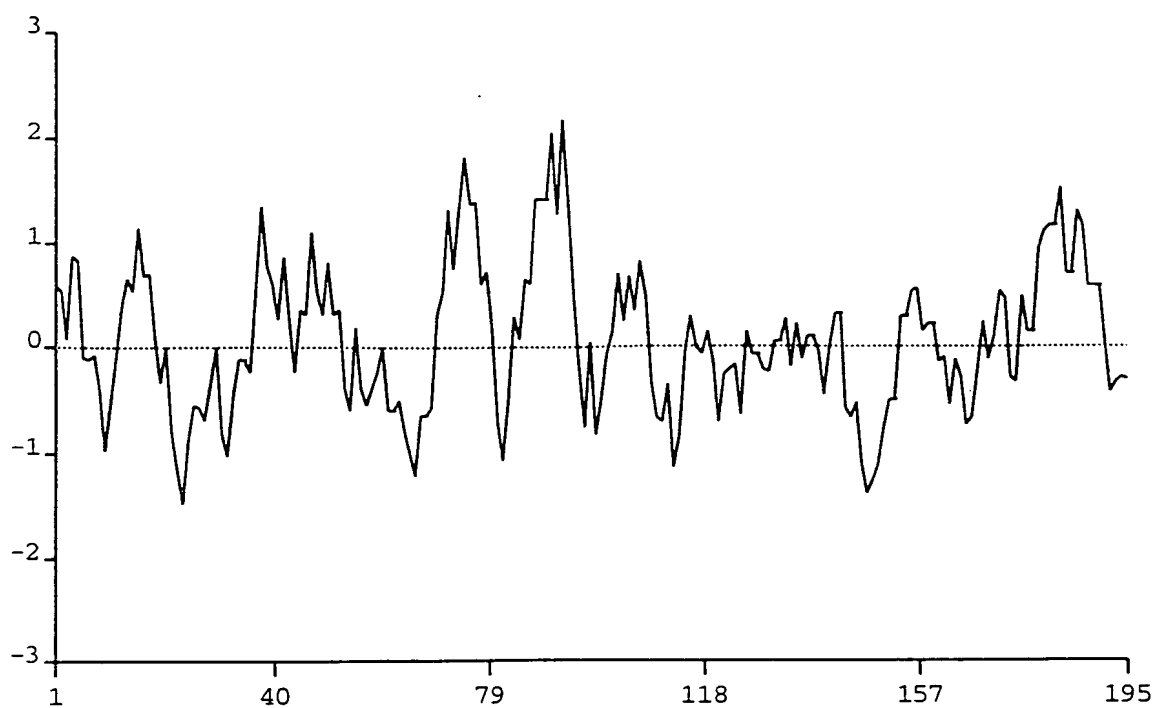


FIGURE 3A

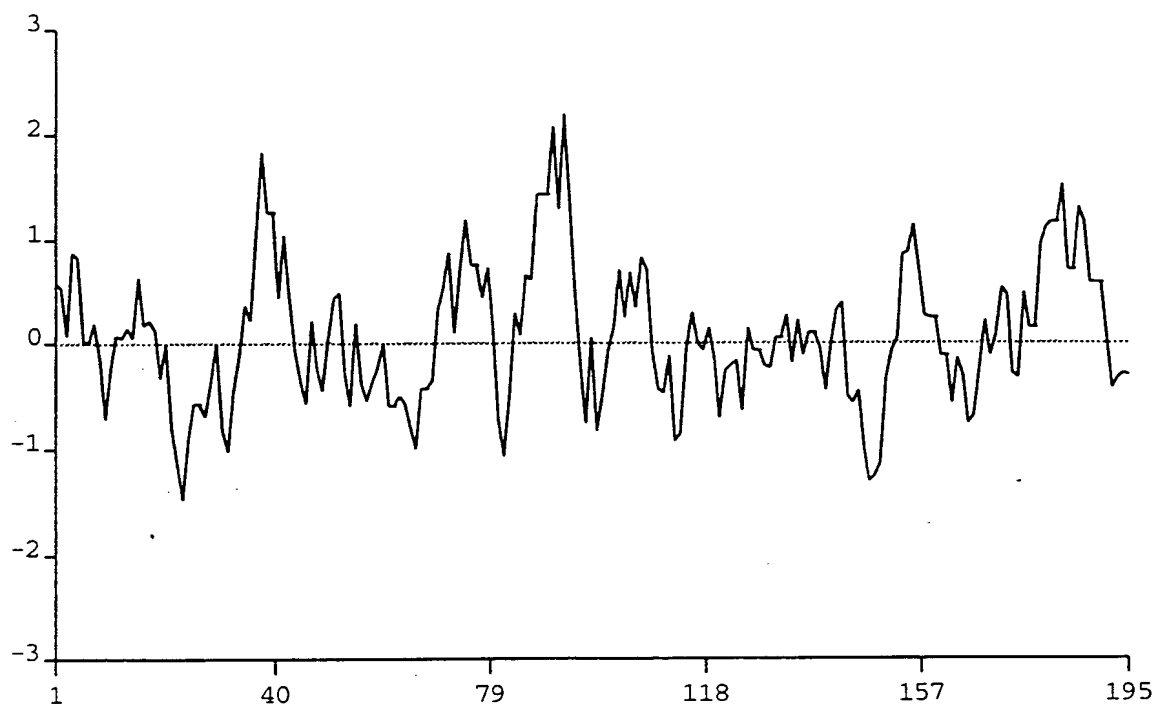


FIGURE 3B